

1 **Sedimentology, faunal content and pollen record of Middle Pleistocene palustrine and lagoonal**
2 **sediments from the Peri-Adriatic basin, Abruzzi, eastern central Italy**

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15

16 **Abstract**

17

18 A 25 m-thick outcrop section exposed at Torre Mucchia, on the sea-cliff north of Ortona, eastern
19 central Italy, comprises a rare mid-Pleistocene succession of shallow-water and paralic sediments along
20 the western Adriatic Sea. An integrated study of the section, including facies and microfacies analyses
21 and characterization of paleobiological associations (mollusks, fishes, ostracods, foraminifers and
22 pollen), enable a detailed reconstruction of the paleoenvironmental and paleoclimatic conditions during
23 deposition. The shallow-water deposits include a transgressive, deepening- and fining-upward
24 shoreface to offshore-transition facies succession overlain by a regressive shoreface-foreshore
25 sandstone body with an erosive base and a rooted and pedogenically altered horizon at the top that
26 imply deposition during sea-level fall. This forced regressive unit is overlain by paralic strata forming a
27 transgressive succession comprising palustrine carbonates and back-barrier lagoonal mudstones. The
28 palustrine carbonates exhibit some of the typical features encountered in palustrine limestones
29 deposited within seasonal freshwater wetlands (marl prairies). Following the sea-level rising trend, the
30 freshwater marshes were abruptly replaced by a barrier-lagoon system that allowed deposition of the

31 overlying mud-rich unit. Within these deposits, the faunal assemblages are consistent with a low-
32 energy brackish environment characterized by a relatively high degree of confinement. The pollen
33 record documents the development of open forest vegetation dominated by *Pinus* and accompanied by
34 a number of mesophilous and thermophilous tree taxa, whose composition supports a tentative
35 correlation with Marine Oxygen Stage 17. The new pollen record from Torre Mucchia improves our
36 understanding of the vegetation development in the Italian Peninsula during the Middle Pleistocene and
37 sheds new light on the role played by the most marked glacial periods in determining the history of tree
38 taxa.

39

40 **Keywords:** Middle Pleistocene, Micropaleontology, Pollen, Mollusks, Fishes, MIS 17, Palustrine
41 carbonates

42

43 **Introduction**

44

45 Nearshore, palustrine, and lagoon systems are very sensitive to variations in sea level and/or sediment
46 availability and their sediments represent ideal targets to assess past and present climate changes (e.g.
47 Rossi et al., 2011; Amorosi et al., 2012, 2013; Sarti et al., 2015). The present contribution illustrates
48 and interprets for the first time the paleontological content of nearshore and paralic strata exposed at
49 the top of the middle Pleistocene Qm2 Unit in eastern central Italy. The studied section is the richest
50 source of paralic mollusk, fish, and pollen assemblages documented so far in the Middle Pleistocene of
51 the western Adriatic Sea and provides a rare opportunity to increase the modest knowledge available on
52 these faunas and floras.

53

54 A 25 m-thick outcrop section of shallow-water and paralic sediments was studied through a
55 multidisciplinary approach involving a detailed description of sedimentary facies and analysis of
56 micropaleontological (planktonic and benthic foraminifers, ostracods and pollen) and
57 macropaleontological (mollusks and fishes) fossil assemblages. Identification of this fauna and flora
58 has significance for understanding the overall paleoecological settings and climatic conditions in which
59 these sediments were deposited.

60

61 **Geological setting**

62

63 The central Apennines are part of an E-NE verging fold-and-thrust belt built up since the Late
64 Oligocene in response to the westward subduction of the Adria micro-plate (i.e., a promontory of the
65 Africa Plate) underneath the European Plate (Malinverno and Ryan, 1986; Ricci Lucchi, 1986;
66 Doglioni, 1991). Progressive tectonic accretion and loading of subducting lithosphere in the Apennine
67 subduction complex (Doglioni, 1991) resulted in the development of a highly articulated foreland basin
68 system at the front of the chain, filled up by relatively thick, diachronous turbidite successions (Ricci
69 Lucchi, 1986). Thrust fronts propagated toward the foreland, involving progressively younger and
70 easternmost foredeep deposits into the orogenic wedge and, at the same time, gradually shifted the
71 depocenters of both foredeep and wedge-top basins further to the east (i.e., Ricci Lucchi, 1986; Ori et
72 al., 1991; Artoni, 2013). The Plio-Pleistocene Peri-Adriatic foredeep records the latest evolutionary
73 stages of the Apennines chain and extends from the Po Plain to the north to the Gulf of Taranto to the
74 south (Fig. 1A). During the latest Miocene to Pleistocene time interval the shape of the central portion
75 of the Peri-Adriatic foredeep, also known as the Pescara depocenter (Ghielmi et al., 2013), was affected
76 by a large variability in space through time, ranging from regular elongated shape to irregular shape,
77 from simple foredeep to fragmented foredeep (Ori et al., 1991). The Quaternary depositional and
78 structural history of this central portion is framed into a large-scale sequence-stratigraphic scheme
79 (Cantalamessa et al., 1986). Specifically, within the traditional subdivision into Quaternary marine
80 (Qm) and Quaternary continental (Qc) unconformity-bounded stratigraphic units, regionally
81 correlatable unconformities divide the Quaternary portion of the basin fill into three major allogroups
82 (Qm1, Qm2, and Qc from oldest to youngest). The unconformities separating these units are well
83 recognizable along the basin margins and document the combined effects of global climate changes and
84 phases of basin reorganization linked to the effects of long-term and regional-scale tectonics (e.g., Ori
85 et al., 1991; Artoni, 2013; Bigi et al., 2013). Overall, these allogroups record a pronounced regressive
86 trend recorded by an upward progression from slope and shelf mudstones of Qm1 (Cantalamessa et al.,
87 2009; Di Celma, 2011; Di Celma et al., 2010, 2013, 2014, 2016a) through littoral sandstones and
88 conglomerates of Qm2 (Cantalamessa and Di Celma, 2004; Di Celma et al., 2016b) to conglomerate-
89 dominated fluvial deposits of Qc (Di Celma et al., 2015). Outcrop equivalents of the Qm and Qc units,
90 displaying strong similarities in terms of both lithology and vertical stacking of facies, have been

91 identified along the Adriatic side of Italy in the Emilia, Marche, and Molise Apennines (Cantalamessa
92 and Di Celma, 2004; Amorosi et al., 2009; Bracone et al., 2012a, 2012b; Gunderson et al., 2014). By
93 early Calabrian times onwards, the thrust-related structures ceased to be active and the central Peri-
94 Adriatic foredeep was uplifted at an average rate of 0.8–1.0 m/ka (Centamore and Nisio, 2003; Pizzi,
95 2003).

96

97 **Study area and methodology**

98

99 Di Celma et al. (2016a) subdivided the sediments of the Middle Pleistocene Qm2 Unit exposed within
100 the Ortona area, eastern central Italy (Fig. 1), into a stack of three unconformity-bounded sequences,
101 namely Qm2₃, Qm2₂, and Qm2₁ from older to younger. Given that this study examines the distal
102 outcrops of the Qm2 Unit, the top-down nomenclature allows potentially older sequences to be
103 identified and consecutively numbered in the extensive exposures of the Qm2 Unit that continues to the
104 west. The present study documents the stratigraphic context, paleontological content and related
105 paleoenvironmental signal of shallow-water and paralic sediments exposed along coastal cliffs to the
106 north of Ortona, where the upper portion of Qm2₂ and the lower portion of the overlying Qm2₁ crop
107 out. The studied sedimentary section is difficult to access because the vertical cliffs are more than 20 m
108 high. Only one locality, called Torre Mucchia (42° 22.677'N - 14° 22.595'E) was found safely
109 accessible (Fig. 2A). The section was measured on a decimeter resolution and the sedimentological
110 characterization of the individual facies associations was focused on the registration of lithology, grain
111 size, primary sedimentary structures, degree and type of bioturbation, paleosols, and the presence of
112 fossils and accessory materials, including roots and plant fragments.

113

114 Field observations and measurements of the cemented chalky limestones have been integrated with
115 sampling of oriented sediment blocks for micromorphological and microfacies analysis. Eight large
116 hand samples were collected and thin sections were prepared from each of them. Thin sections were
117 examined and described following Stoops (2003), Alonso-Zarza and Wright (2010), Freytet and
118 Verrecchia (2002) and Durand et al. (2010). Large gastropod specimens, preserved as molds, were
119 manually collected directly in the field.

120

121 Quantitative analyses on pollen were performed on a total of 29 samples, collected at about 0.2 m
122 intervals along the lower 8 m of the overlying sandy mudstones and prepared according to standard
123 maceration procedures, including treatment with HCl (10%), HF (50%), and NaOH (10%) for 10
124 minutes. Only 12 samples contained suitable pollen concentrations, in a modest state of preservation.
125 The raw counts of pollen analysis are presented as Supplementary Material. The results of pollen
126 analysis are represented as a percentage diagram plotted by means of the program Psimpoll 4.27
127 (Bennett, 2009).

128 Four bulk sediment samples were collected at 2 m intervals for micropalaeontological (ostracods and
129 foraminifers) analyses from the same stratigraphic interval. Approximately 100 g of sediment were
130 picked from each sample, dried at 50°C for 24 h and soaked in water plus H₂O₂ (10%) for 12 h. The
131 samples were subsequently washed under running water through a 63 µm mesh, dried again, and then
132 observed under a binocular microscope.

133

134 The primary source of paleontological remains (ostracods, foraminifers, fishes and mollusks) in the
135 studied succession is a 0.4 m thick, mollusk-dominated fossiliferous interval embedded within the
136 lagoon sediments (Fig. 2A, B). Fish otoliths and mollusks were obtained from the processing of a 20 kg
137 bulk sample amalgamated from multiple replicates (5 kg each) collected from different lateral spots of
138 this rich fossiliferous horizon. The shell-rich sediments were disaggregated in a solution of water and
139 H₂O₂ (10%), sieved through 1 mm mesh sieves with tap water and then dried. The residue from the
140 sample was sorted a few grains at a time, and all otoliths and mollusks were handpicked from the
141 remaining sediment. Overall, both fish otoliths and mollusks are relatively well preserved and fine
142 structures necessary for taxonomic identification are usually clearly recognizable.

143

144 **Results**

145

146 The Middle Pleistocene Qm2 Unit exposed north of Ortona comprises an interval up to 36 m thick of
147 weakly consolidated sandstones, mudstones, conglomerates and minor carbonates, deposited in a
148 variety of continental, paralic, and shallow-marine environments. The detailed sequence stratigraphic
149 interpretation, based on key surfaces and stratal units that are traceable across the entire study area, has
150 been presented elsewhere (Di Celma et al., 2016a) and is not reiterated here. The description and

151 interpretation of the five facies associations documented in the studied section are provided below,
152 whereas a view of their vertical stacking pattern is shown in Figure 2C.

153

154 *Facies associations*

155 In the 25 m-thick studied succession (upper portion of Qm2₂ and lower portion of Qm2₁), the
156 sedimentological analyses led to distinguish five major facies associations. In ascending order, these
157 facies associations are described below in terms of sedimentological and paleontological features.

158 Figure 2C shows a synthetic interpretation of the outcrop section.

159

160 *Lower shoreface bioturbated massive sandstone (LSh)*

161 This facies association is 5 m thick and consists of thoroughly bioturbated, fine- to medium-grained
162 silty sandstones. These sediments are generally structureless, although poorly preserved mm-scale
163 plane-parallel and low-angle laminations can locally be identified.

164 Fragments of marine mollusks and rare well-rounded extraformational clasts averaging 1-1.5 cm in
165 diameter occur randomly scattered throughout the facies association. Discrete intervals of soft
166 sediment-deformed sandstone beds, as much as 1.5 m thick and characterized by recumbent folds,
167 occur intermittently along the outcrop belt, more often at the top of this facies association and close to
168 its vertical transition with the overlying offshore-transition facies association.

169

170 The ichnofossil assemblage is largely dominated by unusually large, densely packed, irregularly
171 meandering, horizontal to gently inclined, cylindrical to subcylindrical, unbranched burrows of
172 *Macaronichnus* sp., but also comprises some isolated burrows of *Rosselia socialis*, a vertically-oriented
173 spindle- to funnel-shaped, mud-lined tube, and *Skolithos linearis*.

174

175 The overall paucity of physical sedimentary structures, restricted to the presence of rare plane-parallel
176 and low-angle laminations, the absence of storm-induced sedimentary structures, such as hummocky
177 and swaley cross-stratification, and the pervasively burrowed fabric, indicate that storm waves and
178 currents did not deposit beds significantly thick enough to shield them from total reworking by
179 burrowing organisms and that low-energy conditions prevailed during deposition of this facies
180 association. Unstable sea-floor conditions are indicated by the occurrence of soft-sediment structures.

181 Triggering mechanisms are unknown, but this instability may have been imparted by seismic
182 disturbances or, less likely, by infrequent large magnitude storms.

183

184 The *Macaronichnus* burrows record intrastratal trails produced by infaunal detritus-feeding polychaete
185 worms. With the exception of their unusually large size, the Torre Mucchia *Macaronichnus* burrows
186 are morphologically very similar to those of standard (small) *Macaronichnus segregatis* described by
187 Clifton and Thompson (1978). Similar large *Macaronichnus* burrows have been reported from both
188 ancient (Nara, 2002; Aguirre et al., 2010) and modern (Seike et al., 2011) shoreface deposits. Trace-
189 fossil assemblages overwhelmingly dominated by robust deposit-feeding structures, such as large
190 *Macaronichnus* and *Rosselia*, and with subordinate traces of suspension-feeding organisms, such as
191 *Skolithos*, corresponds to a proximal expression of the *Cruziana* Ichnofacies (Pemberton et al., 2012
192 and references therein). Both the sedimentary features and the trace fossils of this facies association
193 indicate that deposition occurred above fair-weather wave base, in a well oxygenated, low-energy
194 lower-shoreface setting with infrequent storm influence.

195

196 *Offshore-transition bioturbated mudstones and fine grained sandstones (OfTr)*

197 At Torre Mucchia this facies association is a 5 m-thick, thoroughly burrowed mixture of sandstone and
198 mudstone. Sandstones are fine-grained and generally preserved in rare, sharp based layers that range
199 from mm- to cm-scale thicknesses and show hints of some horizontal planar to low-angle laminations.
200 Mudstones are characterized by bed-scale variations from moderately to thoroughly bioturbated silty
201 mudstones. Pervasive bioturbation, however, is very common and obscures most of remnant physical
202 sedimentary structures, rendering an overall churned fabric to this facies association. Mollusk shells are
203 present but sparse and consist primarily of disarticulated bivalves found either dispersed throughout
204 this facies or as small clumps. Apart from some little clumps of *Ostrea edulis*, characterizing faunal
205 elements at the outcrop level are the infaunal species *Panopea glycymeris*, mainly preserved as
206 conjoined individuals in vertical (life) or weak oblique position, and *Glycymeris nummaria* (= *violacescens*)
207 preserved both as single and articulated valves. Sparse remains of *Pinna* sp. consist of
208 single, very fragmented valves.

209

210 This facies association preserves a record of alternating high- and low-energy conditions. The
211 burrowed mudstones reflect periods of fair-weather conditions and deposition of mud from suspension.

212 The interbedded sharp-based, horizontal planar to low-angle laminated sandstone beds suggest
213 deposition during episodic storms, when sand was eroded from the upper part of the shoreface and
214 transported basinward. Due to their limited thickness and the intense reworking by burrowing
215 organisms, the preservation potential of these sandstone beds was low.

216

217 The macrofossil fauna documented within this facies association is composed of still living species (*P.*
218 *glycymeris*, *G. nummaria*, and *Pinna* sp.) that, today, are characteristic inhabitants of inner to middle
219 shelf muddy soft-grounds (Marasti, 1991; Bernasconi and Robba, 1993). Sediment grain size, complete
220 biogenic homogenisation, and macrofossil content converge to indicate that this mudstone-rich facies
221 association has been deposited between fair-weather and storm-wave base and reflects accumulation in
222 an offshore-transition depositional environment. The high degree of bioturbation observed in this facies
223 association indicates that storm events were likely followed by long periods of low-energy conditions
224 favoring the intense reworking of sediments.

225

226 *Upper shoreface to foreshore cross- and plane-parallel laminated sandstones (UShFo)*

227 This facies association is in the upper part of Qm2₂ and is about 6 m-thick, composed of medium- to
228 coarse-grained sandstone beds with swaley cross-stratification in the lower part, plane-parallel
229 laminated and high-angle trough and planar cross-bedded sandstones in sets several tens of centimeters
230 thick in the middle part, and plane-parallel laminated sandstones in wedge-shaped sets dipping seaward
231 at low angles in the uppermost part. Due to intense bioturbation by rootlets, the uppermost 1.5 m of this
232 facies association is usually massive, with the root traces subtending from a laterally extensive, planar
233 top surface associated with a reddish palaeosol (Fig. 2D). Palaeocurrent measurements in high-angle
234 planar cross-beds indicate a dominant southeastward flow direction. Local patches displaying intense
235 reworking by large *Macaronichnus*, with horizontal burrows generally running parallel to the bedding
236 planes account for most of the bioturbation in this facies association.

237 The lower boundary of this sandstone package is sharp, typically flat and erosional onto the underlying
238 mud-rich sediments (Fig. 2C), whereas the upper boundary is a surface of subaerial exposure, typically
239 marked by a rooted, reddish paleosol (Fig. 2D). About 5 km southeast of the Torre Mucchia Section,
240 well preserved within this subaerially weathered horizon, Agostini et al. (2007) documented a

241 vertebrate fossil fauna including the hippopotamus *Hippopotamus* ex gr. *H. antiquus*, the straight-
242 tusked elephant *Elephas (Loxodon) antiquus*, and the red deer *Cervus elaphus*.

243

244 Swaley cross-stratification (Leckie and Walker, 1982) is usually associated to the combination of
245 powerful storm-generated unidirectional and oscillatory flows at and above the limit of the fair-weather
246 wave base (e.g. Dott and Bourgeois, 1982; Arnott, 1992). Planar and trough cross-stratification denotes
247 migration of two- and three-dimensional dunes. The abundance of shoreline-parallel paleocurrents
248 implies that strong longshore currents transported large volumes of sand (e.g., Hart and Plint, 1989;
249 Walker and Plint, 1992; Frébourg et al., 2012). The relatively low intensity of bioturbation and
250 impoverished ichnological diversity together with the preserved sedimentary structures are indicative
251 of deposition in a storm/wave influenced shoreface environment, in which high storm-wave frequency
252 and energy resulted in amalgamated sandstone beds (e.g., Bourgeois and Leithold, 1984; Clifton and
253 Dinger, 1984; Church and Middleton, 2003). In this setting, both two- and three-dimensional dunes
254 migrated episodically, as evidenced by the local occurrence of completely bioturbated bedding planes,
255 which represent periods of colonization of a shifting sandy substrate by polychaete worms. The
256 seaward-dipping, low-angle, plane-parallel laminated sandstones in the uppermost part of this facies
257 association are interpreted as foreshore deposits where plane-parallel-lamination formed under upper
258 flow regime due to intense swash and backwash processes. This interpretation is supported by the
259 occurrence of root traces and a capping paleosol representing a period of sub-aerial exposure and
260 pedogenic alteration of the deposits. This sharp-based, regressive succession indicates limited
261 accommodation in front of the prograding shoreface and represents a relative drop in sea level.
262 Accordingly, it is placed in the falling stage systems tract and its erosional base is interpreted as a
263 regressive surface of marine erosion.

264

265 *Palustrine chalky carbonates (Pal)*

266 This facies association, up to 0.5 m in thickness, characterizes the lowermost portion of Qm2₁ and
267 shows an unconformable lower boundary with the underlying upper shoreface-foreshore sandstones,
268 highlighted by the occurrence of a rooted red paleosol (Fig. 2D). The facies association consists of
269 chalky carbonates and includes three main facies, namely A, B, and C (Fig. 3A), composed of

270 intensively rooted wackestone, packstone, grainstone and mudstone beds, rich in gastropods,
271 intraclasts, calcareous and siliciclastic sand.

272

273 The basal facies A (Fig. 3B) is composed of fine to medium packed sands (7.5YR 8/3) with abundant
274 blackish to reddish Fe/Mn precipitation features along irregular wavy sub-horizontal laminae and by
275 reddish to orange (5YR 4/2) sub-vertical features that cross the laminated sands (Fig. 3B). Macro- and
276 micromorphological observations do not reveal any carbonate leaching or clay illuviation features. The
277 basal surface of this facies is also characterized by the presence of strongly cemented, cylindrical root
278 traces (Fig. 3C). Two main microfacies can be recognized, namely A-1 and A-2. The basal A-1
279 microfacies (Fig. 4A) is made up of packed sands (c/f ratio 1), with light micritic groundmass and very
280 abundant blackish impregnation features scattered or roughly aligned along undulated, irregular sub-
281 horizontal surfaces. The coarse fraction is composed of predominant subangular to angular chert and
282 quartz grains with subordinated rounded to subrounded limestone grains. An abrupt change marks the
283 appearance of the upper A-2 microfacies. The coarse fraction decreases (c/f ratio 1/10), the groundmass
284 is darker and made of cyanobacterial clotted micrite (Fig. 4B). Fe/Mn features concentrate along voids,
285 cracks or diffused in the groundmass with dendrolithic forms (Fig. 4C). Microsparitic or sparitic
286 features are rare and mainly related to dissolution of shells or as coatings of small biological voids.
287 A clear undulated boundary marks the transition to the overlying facies B (Fig. 3A), which is
288 characterized by a weak lamination, abundant presence of gastropods, common fine and thin vertical
289 tubules, abundant yellowish-orange to reddish- black Fe/Mn masses, nodules or concentrations within
290 the vertical tubules. A single microfacies is recognized in facies B, with a c/f ratio 1/4 and decreasing
291 Fe/Mn features toward the top. These are coatings and infillings of planes, voids and cracks (Fig. 4D),
292 locally with precipitation of hematite, and quasi-coatings around voids and impregnative masses within
293 the groundmass. The groundmass is dominated by cyanobacterial clotted dark micrite irregularly
294 alternated with lighter micrite. Microsparite and sparite are rare, related to dissolution of shells or as
295 coatings on rounded biological voids (Fig. 4E).

296

297 A diffused boundary separates facies B from the overlying muddy facies C (Fig. 3A). The color of
298 facies C is lighter (10YR 8/2), thin vertical root traces filled with yellowish orange silts are commons
299 (Fig. 3D); Fe/Mn features are scarce whereas large CaCO₃ nodules or irregular concentrations are

300 common and gastropods are abundant (Fig. 3D) and include a mixture of terrestrial pulmonates and
301 freshwater prosobranchs (*Viviparus* sp., *Cepaea* sp., *Valvata* sp., and *Bithynia* sp.; D. Esu, 2014 and M.
302 Magnatti, 2016 personal communications). Microfacies C-1 is observed at the base. This is dominated
303 by irregularly distributed dark cyanobacterial clotted micrite (Fig. 4F) with very abundant shells, shell
304 fragments and foraminifera. The coarse mineral fraction is subordinated (c/f ratio 1/8). Orange to
305 reddish Fe precipitation features are concentrated within planes, cracks and biological voids (Fig. 3E).
306 Microsparitic and sparitic features are very rare, fine and mostly related to dissolution of shells.
307 Toward the top the microfacies progressively changes into the Microfacies C-2, characterized by
308 irregularly alternating dark cyanobacterial clotted micrite and light micrite dominated groundmass (Fig.
309 4G). The darker cyanobacterial micrite is also characterized by abundant microsparitic and sparitic
310 fillings and coatings along cracks and biological voids (Fig. 4H). Fe/Mn features are scarce and made
311 of small nodules and impregnative features that are more abundant within the darker groundmass
312 patches.

313

314 This facies association represents marl prairie palustrine carbonates precipitated both biogenically and
315 physico-chemically within very shallow carbonate marshes in seasonal freshwater wetlands (e.g. Platt
316 and Wright, 1992; Alonso-Zarza et al., 2006; MacNeil and Jones, 2006; Reuter et al., 2009) that were
317 characterized by slowly moving or stagnant water. Macro- and micromorphological observations
318 indicate that the rooted paleosoil that developed on top of the marine sands represents a short-lived
319 event of exposure, lacking any evidence of weathering of the mineral fraction, carbonate leaching and
320 clay neoformation or illuviation. The main observed processes are related to a vadose environment,
321 typically cyanobacterial clotted micrite alternated with light micrite. The rare microsparitic coatings
322 and fillings suggests very short-lived periods of complete subaerial exposure. However, phreatic
323 cements are very rare, more abundant in the upper facies suggesting the rising of the water table and
324 the possible flooding of the environment. Water table oscillation in vadose environment is also
325 indicated by the abundance of Fe/Mn features (marmorization, *sensu* Freytet and Verrecchia, 2002).

326

327 *Lagoon interbedded grey-green mudstones and very fine-grained sandstones (Lg)*

328 This facies association forms the bulk of Qm2₁ and is between 15 m and 18 m thick, of which only the
329 basal 8 m are continuously exposed in the studied section. This sediment package consists of organic-

330 rich (essentially plant debris), grey to green massive mudstones interbedded with subordinate,
331 decimetric- to centimetric-thick, very fine-grained sandstone beds showing no overall grain-size trends.
332 Traces of bioturbation and scattered mollusk shells are found throughout the sampled interval.

333

334 A distinctive, 0.4 m-thick, highly-fossiliferous horizon occurs at the top of the exposed section and
335 represents the richest source of paleontological data, yielding abundant mollusks, fishes, benthic
336 meiofauna, and planktonic foraminifers. The mollusk assemblage consists of the macromollusk
337 *Cerastoderma glaucum* (mainly represented by juvenile individuals) and *Abra segmentum* along with a
338 slightly more diverse micromollusk group including extremely abundant hydrobiid mudsnails, *Bittium*
339 *scabrum*, *Planorbarius corneus*, *Rissoa membranacea*, *Theodoxus fluviatilis*, *Tragula fenestrata* and
340 *Valvata piscinalis* (Fig. 5). Most of the faunal components are relatively well preserved and a minor
341 amount of shells show a poor state of preservation.

342

343 The vertebrate assemblage is composed of fish remains, including saccular otoliths and a few teeth
344 (Fig. 6). The fish otoliths are scarcely diversified, revealing a remarkable oligotypic character of the
345 fish assemblage. About 75% of the recognized specimens belong to the big-scale sand smelt *Atherina*
346 *hepsetus*. The remaining part of the assemblage is represented by gobies morphologically very similar
347 to those pertaining to the *Pomatoschistus minutus* species complex (see Webb, 1980; Nolf, 2013).

348 The ostracod fauna exhibits the overwhelming dominance of the highly euryhaline *Cyprideis torosa*
349 and a few scattered valves of *Leptocythere* sp. accompanied by a benthic foraminiferal fauna
350 characterized by an oligotypic assemblage dominated by the euryaline species *Ammonia beccarii*
351 (about 80%) and subordinately by *Haynesina paucilocula* (10%), *Valvulineria perlucida* (7%) and
352 *Cribrononion granosum* (3%) (Fig. 7). The samples contain mostly reworked benthic and planktonic
353 foraminifers, usually poorly preserved, with eroded and broken shells. Reworked benthic foraminifers
354 belong to the following taxa: *Hyalinea baltica*, *Elphidium crispum* and *Cibicides* sp., whereas the
355 planktonic foraminifers, usually very small in size, are *Globorotalia inflata*, *Globigerinoides* sp., and
356 *Globorotalia* sp.

357

358 The rich organic fraction suggests deposition in a low-energy, restricted, brackish-water, lagoonal
359 environment occasionally affected by high-energy events represented by back-barrier overwash
360 sandstone beds.

361

362 Monospecific ostracod assemblages dominated by *C. torosa* are interpreted to represent environmental
363 stress. This species is typical of brackish-water and lagoonal environments (Athersuch et al., 1989;
364 Meisch, 2000) and it is also highly euryhaline, showing an adaptability to salinities from 0.4 to 150‰
365 (Neale, 1988) and therefore an opportunistic behavior (Frenzel and Boomer, 2005). The comparison
366 between the abundance of the species *A. beccarii* with those of the species *H. paucilocula*, *V. perlucida*
367 and *C. granosum* suggests the presence of distinct lagoonal environments (Albani and Serandri-
368 Barbero 1982; Serandrei-Barbero et al., 2005, 2006; Zecchin et al., 2009). In this case, a high
369 percentage of *A. beccarii* and a low abundance of *H. paucilocula*, *V. perlucida* and *C. granosum*
370 identify an inner lagoonal environment with hyposaline conditions (Albani and Serandrei Barbero,
371 1982; Donnici et al., 1997; Serandrei-Barbero et al., 2005, 2006).

372 The mollusk record is characterized by some brackish-water taxa (among which Hydrobiidae, *A.*
373 *segmentum* and *C. glaucum*, in order of abundance) typical of the euryhaline and eurythermal lagoons
374 (LEE *sensu* Peres and Picard, 1964). Hydrobiidae are herbivores or surface detritivores burrowers
375 adapted to euryhaline conditions ranging from 2 to 34‰ even if usually they prefer the range 6-25‰
376 (Fretter and Graham, 1978), *C. glaucum* is a filter-feeder normally living in water bodies characterized
377 by salinities between 18 and 37‰ (Vatova, 1981), and *A. segmentum* is a browser-burrower which
378 thrives in a salinity range of 14-27‰. Therefore the contemporaneous occurrence of these three taxa
379 might suggest a range of salinity between 18 and 27‰. However, the presence of *P. corneus*, *V.*
380 *piscinalis*, a fresh-water species which tolerates slightly brackish waters (Grigorovich et al., 2005), and
381 of *R. membranosa*, which regularly occurs at salinities of 7–10‰ (Waren, 1996), along with the
382 absence of marine affinity taxa point toward mesohaline waters far from marine influence. This is not
383 in contrast with the presence of *A. segmentum* and juvenile individuals of *C. glaucum* because these
384 taxa may tolerate low salinities as low as 5‰ (Di Rita et al., 2011), a value considered as a critical
385 threshold for distribution of the lagoonal taxa (Cognetti and Maltagliati, 2000). Data from the mollusk
386 content are consistent with a low-energy brackish environment with very limited marine influence,
387 such as an inner lagoon, in which the presence of algal/plant meadows is suggested by the occurrence

388 of browsers. According to the “confinement model” proposed by Guelorget and Perthuisot (1992) the
389 paralic mollusk assemblage may be referable to a lagoon environment of zone IV corresponding to a
390 rather high degree of confinement. Such a zonal reference is supported also by the sketch proposed by
391 Breber et al. (2000) solely based on bivalves.

392

393 Sand smelts and gobies are common constituents of paralic assemblages in terms of relative abundance,
394 representing among the best adapted fishes for this type of habitats (e.g., Harrison and Whitfield,
395 1995). Sand smelts are small-sized pelagic planktivores that usually reside in paralic biotopes in which
396 they can complete their short life cycles (e.g., Nordlie, 2003; Franco et al., 2006). More particularly,
397 *Atherina hepsetus* exhibits a strong tolerance to broad salinity variations, inhabiting waters with
398 salinities ranging from 1.6 to 4.5‰ (Kiener and Spillmann, 1969). Gobies of the genus *Pomatoschistus*
399 are very abundant in shallow coastal areas, estuaries, coastal lagoons and marine bays of the
400 Mediterranean and Eastern Atlantic Ocean where they feed on small benthic invertebrates, pelagic
401 copepoda and small fishes (e.g., Hamerlynck and Cattrijsse, 1994). From an ecological point of view,
402 all the recognized fish taxa can be assigned to the guild of estuarine residents (see Elliot and Dewailly,
403 1995), which includes species that spend their life in brackish biotopes influenced by marine waters.
404 Overall, the exclusive presence of taxa belonging to this guild is indicative of soft bottoms and densely
405 vegetated biotopes (Paterson and Whitfield, 2000). The absence of marine and/or freshwater migratory
406 species seems to be indicative of the reduced influence of both the marine inflow and freshwater
407 outflow, suggesting that the original depositional environment was characterized by a relatively high
408 degree of confinement (Zones III and IV of the confinement scale of Guelorget and Perthuisot, 1992).

409

410 *Pollen analysis*

411 The pollen record of the Torre Mucchia section shows the development of a forest vegetation
412 dominated by *Pinus* (>40%), accompanied by a number of mesophilous and thermophilous tree taxa,
413 including deciduous and evergreen *Quercus*, reaching 13 and 9%, respectively, *Alnus* (8%), *Carpinus*
414 *orientalis* type (7%), *Carpinus betulus* (2%), *Picea* (7%), *Juniperus* (7%), *Abies* (6%), and several
415 other tree taxa in low percentages (Fig. 8). A significant presence of *Tsuga* pollen, attaining 9% in one
416 sample, *Carya*, reaching values of 5% in the bottom samples, and *Zelkova* (12%), accompanied by
417 scattered presence of *Pterocarya*, are of special interest, as these tree taxa disappeared from Europe in

418 the course of the Middle Pleistocene (Bertini, 2010; Magri and Palombo, 2013). Chenopodiaceae (max.
419 30%), *Artemisia* (17%), Cichorioideae (15%), Asteroideae (6%), Poaceae (13%), and Apiaceae (7%)
420 are common herbaceous taxa throughout the record.

421

422 As a whole, the pollen record represents an open forest, whose floristic composition clearly reflects a
423 mixture of both mountain (Majella Massif) and coastal (Adriatic Sea) environments and suggests that a
424 variety of vegetation belts were present in the surrounding areas: i) a mountain belt that includes high-
425 altitude, water-demanding trees, such as *Picea*, *Abies*, *Tsuga*, and *Betula*; ii) a mid-altitude belt
426 including *Carya*, *Carpinus*, deciduous *Quercus*, *Alnus*, and *Corylus*; and iii) a warm coastal belt
427 including evergreen *Quercus* and possibly *Zelkova*. The percentage values of Asteraceae (between 6%
428 and 32%) and Apiaceae (up to 7%) indicate the presence of wide open areas dominated by steppe
429 elements. Together with the abundance of *Pinus*, they suggest that the recorded environments do not
430 represent the fully developed forest conditions that correspond to the warmest and wettest phases of
431 interglacial periods, which are generally dominated by broadleaved trees. The continuous and
432 significant frequencies of chenopods suggest saline conditions linked to the proximity of the sea, as
433 also indicated by the presence of foraminiferal linings and dinocysts. Similar conditions were recorded
434 in other pollen diagrams from coastal environments, in connection with inputs of marine water into
435 coastal lakes (Di Rita and Melis, 2013; Melis et al., 2015).

436

437 **Discussion**

438

439 *Stratigraphic evolution*

440 In the studied section, the vertical succession of facies associations can be interpreted as related to
441 relative sea-level oscillations and comprises the two youngest sequences exposed in the Ortona area,
442 namely Qm2₂ and Qm2₁ (Fig. 2C). The lower sequence (Qm2₂) includes a transgressive, deepening-
443 and fining-upward shoreface to offshore-transition facies succession overlain by a shoreface-foreshore
444 body with an erosive base and a rooted pedogenic horizon at the top (Di Celma et al., 2016a). The
445 presence at the base of the shoreface-foreshore sandstones of a sharp and scoured surface that truncates
446 to some degree the underlying offshore-transition sediments and the evidence of subaerial exposure and
447 pedogenic modification at its top imply forced regression and deposition during sea-level fall (Plint and

448 Nummedal, 2000 and references therein). The degree of development of paleosols is function of the
449 soil-forming factors that in sedimentary record i.e., regressive-transgressive cycles, can be summarized
450 into time (length of subaerial exposure of the surface; Kraus, 1999) and climate conditions (Interglacial
451 vs. Glacial; Coltorti and Pieruccini, 2006). The weakly developed interfluvial palaeosol described here
452 can be traced laterally into the basal erosional surface of an incised valley fill (Di Celma et al., 2016a)
453 and, therefore, it is interpreted to represent a sequence-bounding subaerial unconformity. In forced
454 regressive stages subaerial unconformities are highly diachronous surfaces that form initially in updip
455 areas and gradually expand in a basinward direction as the sea-level falls and the shoreline regresses
456 (e.g., Plint et al., 2001). The paleosols development continues during subsequent lowstand normal
457 regression and transgression and does not cease until the exposure surface becomes buried by renewed
458 deposition. These observations imply that the degree of development of sequence-bounding interfluvial
459 paleosols is a function of their position along the depositional profile, with the most mature paleosols
460 predicted in the most updip areas and the least developed paleosols expected to form in the most
461 downdip areas, where they have less time to develop. As a result, the poorly developed character of the
462 paleosol exposed at Torre Mucchia can be interpreted as reflecting a distal position of the study section
463 along the subaerially exposed shelf. Moreover, the described paleosol formed during an eustatic forced
464 regression and, therefore, not in fully Interglacial conditions, when long lasting warm and wet climatic
465 conditions favor the formation of strongly developed paleosols as observed also in the nearby areas
466 (Coltorti and Pieruccini, 2006; Di Celma et al., 2015). These would not only explain the low degree of
467 pedological development of the paleosol, but also the abundance of *Pinus* and herbs in the pollen
468 record from the lagoon mudstones, reflecting the initial transgression during a glacial-interglacial
469 transition, as also recorded in other lagoon sites (Di Rita et al., 2015). High percentage values of *Pinus*,
470 generally >30%, accompanied by deciduous trees not exceeding 10%, characterized also the
471 interglacial vegetation of the Cesi section, an early Middle Pleistocene pollen site in central Italy
472 (Bertini, 2000). During the subsequent transgression, the landward migration of a barrier-lagoon
473 system developed behind the transgressing shoreline resulted in the inundation of the previous coastal
474 plain and deposition of palustrine carbonates in a freshwater coastal wetland that eventually evolved
475 into a brackish lagoon within which mud-rich sediments were deposited.

476

477 *Chronological constraints*

478 The compositional correlation between the pollen assemblage retrieved from the lagoon mudstones of
479 the Qm2₁ sequence and other floristic records documented in central-southern Italy, between 42° and
480 39° latitude N (Figs. 1A, 9), provides some additional chronological constraint for the studied
481 succession. *Tsuga*, reaching the significant percentage of 9% at Torre Mucchia, was found in
482 appreciable amounts (>5%) until MIS 18 at Montalbano Jonico, a sequence dated by means of two
483 ⁴⁰Ar/³⁹Ar ages on volcanoclastic layers (⁴⁰Ar/³⁹Ar 773.9 ± 1.3 and 801.2 ± 19.5 ka; Bertini et al., 2015).
484 It was also continuously found at Valle di Manche before and soon after the Matuyama–Brunhes
485 boundary (Capraro et al., 2005). After that time, *Tsuga* was found in two levels (<2.5%) in the deposit
486 of Sessano, correlated with MIS 13 on the basis of a ⁴⁰Ar/³⁹Ar age (437.9 ± 1.9 ka; Russo Ermolli et al.,
487 2010). During the same interglacial, there are only two occurrences of single grains at Vallo di Diano
488 (Russo Ermolli, 1994). Other pollen records from southern Italy correlated with MIS 13 (Acerno,
489 Boiano and Mercure; Russo Ermolli et al., 2015) do not record any pollen grain of *Tsuga* and indicate
490 that its range was markedly fragmented, as it is expected for a tree population that is undergoing
491 extinction. On the whole, the abundance of *Tsuga* in the Qm2₁ sequence suggests an age older than
492 MIS 16 (Fig. 9).

493

494 Another tree taxon that is no longer present in Italy is *Carya*, which is very abundant (>25%) in central
495 Italy during the Early Pleistocene interglacials between 1.8 and 1.2 Ma (Magri and Palombo, 2013).
496 Percentages <10% have been documented at Fontana Ranuccio, in sediments underlying the
497 Matuyama–Brunhes boundary and ascribed to MIS 21 (Corrado and Magri, 2011). During MIS 19, at
498 Montalbano Jonico (Bertini et al., 2015) and Valle di Manche (Capraro et al., 2005) *Carya* never
499 reached values >5%. *Carya* was absent from Vallo di Diano during MIS 15, but continuously present
500 in very low abundance during MIS 13 (Russo Ermolli, 1994). Pollen associated to a macrofossil find
501 dated 530 ka at Carsoli (Sadori et al., 2010) and pollen grains (always <1.5%) at Ceprano (Manzi et al.,
502 2010) confirm its presence in Central Italy during MIS 13. The very last appearances of *Carya* are
503 documented at Boiano, southern Italy, within MIS 9 sediments (Orain et al., 2013). The pollen record
504 from the Qm2₁ sequence, showing maximum values of *Carya* around 5%, fits the progressive reduction
505 of this taxon during the Middle Pleistocene, further supporting an age older than MIS 16 for the Torre
506 Mucchia pollen record (Fig. 9).

507 On the whole, the pollen diagram from the Middle Pleistocene lagoon mudstones of the Qm2₁ sequence
508 is similar to the records correlated with MIS 21–17, while it is clearly distinct from the records
509 corresponding to MIS 15–13 (Fig. 9). This attribution is also supported by the presence of *Pterocarya*,
510 whose appearances become sporadic in the records corresponding to MIS 15–13 (Fig. 9). In contrast,
511 *Carpinus* and *Fagus*, showing low percentages in all the Italian records between 1.5 and 0.7 Ma (Magri
512 and Palombo, 2013), increase to values of 5–10%, in correspondence with high percentages of *Abies*,
513 during MIS 15 and 13 (Fig. 9). The difference in vegetation between MIS 21–17 and MIS 15–13 is also
514 marked by the reduction of *Tsuga* and *Carya*. This turnover in the composition of forest vegetation
515 may be explained by the extreme climatic conditions of MIS 16 (Tzedakis et al., 2006).

516

517 The correlation of the mammal assemblage from the immediately underlying paleosol to an interval
518 bracketed between MIS 19 and MIS 17 (Mazza and Bertini, 2013), based on the paleomagnetic reversal
519 assigned to the Matuyama-Bruhnes transition in the nearby section of Ortona (Agostini et al., 2007),
520 provides an additional chronological constraint, as it implies an age younger than MIS 19. Thus, it is
521 feasible that the studied portion of the Qm2₁ sequence was laid down during the early transgression of
522 MIS 17, not in fully interglacial conditions yet.

523

524 **Conclusions**

525

526 The multi-disciplinary analysis carried out on the Torre Mucchia section, Ortona, integrates
527 sedimentological features with data obtained from the analysis of fish, mollusks, pollen, ostracods and
528 foraminifers found within the sedimentary section. These data provide a rare opportunity to reconstruct
529 the articulate evolutionary scenario of a coastal area along the western side of the Adriatic Sea during
530 Middle Pleistocene and a constraint for the age of the upper part of the Qm2 Unit (Qm2₁ sequence).

531 The stratigraphic unconformity at the top of the forced regressive upper shoreface-foreshore
532 sandstones, marked by a rooted reddish paleosol, records a phase of non-deposition and pedogenic
533 modification in the study area. Above this emerged surface, the palustrine carbonates represent the first
534 record of transgressive deposition and the transgressive surface is located at its lower boundary, at the
535 passage with the underlying rooted paleosol. Sedimentological, paleontological and
536 micromorphological features suggest that these carbonates were deposited within very shallow,

537 seasonal freshwater wetlands probably developed not far from the coeval coastline. As relative sea-
538 level rise and transgression proceeded, the freshwater wetlands evolved into a brackish lagoon in which
539 mud-rich sediments were deposited. The paleontological content and composition of the pollen
540 assemblage provide useful information about the palaeoenvironmental and climatic conditions along
541 the western Adriatic Sea during deposition of the Qm2₁ sequence and a valuable clue to the
542 chronostratigraphy and geochronology of the studied section. Within these deposits, the faunal
543 assemblages suggest that the original depositional environment was characterized by brackish waters
544 and a relatively high degree of confinement. The pollen record documents the development of a forest
545 vegetation dominated by *Pinus* and accompanied by a number of mesophilous and thermophilous tree
546 taxa and significant amounts of herbaceous pollen pointing to the presence of wide open areas. These
547 vegetation features, considered in the light of the correlation with the mammal fauna and the
548 paleomagnetism of a nearby sequence from Ortona (Agostini et al., 2007; Mazza and Bertini, 2013),
549 are suggestive of deposition during an early phase of MIS 17.

550
551 The new pollen record from Torre Mucchia fills a gap of information on the vegetation history in
552 Central Italy, thus significantly improving our understanding of the vegetation development during the
553 Middle Pleistocene and especially of the role played by the most marked glacial periods in determining
554 the reduction or disappearance of tree taxa that are nowadays extinct in the Italian Peninsula.

555
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557
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563
564 **Figure captions**

565

566 Figure 1. (A) Location map of Ortona and the main Middle Pleistocene Italian sites (between ca. 39°
567 and 42° N) cited in the text for their palynological data. (B) Enlarged satellite image showing
568 location of the study area.

569 Figure 2. (A) Panoramic view of the Torre Muchia section. Dashed and dotted lines as indicated in C.
570 (B) Detail of the shell-rich interval embedded in the lagoon mudstones. (C) Vertical distribution of
571 facies associations and sequence stratigraphic interpretation of the studied section. The key stratal
572 surfaces are highlighted by dashed and dotted lines. LSh, lower shoreface; OfTr, offshore-
573 transition; UShFo, upper shoreface-foreshore; Lg, lagoon mudstones. (D) View of outcrop
574 showing the vertical transition from sequence Qm2₂ to sequence Qm2₁. UshFo, upper shoreface-
575 foreshore sandstones; Ps, paleosol; Pal, palustrine carbonates; Lg, lagoon mudstones.

576 Figure 3. Compilation of field photographs showing the main sedimentological characteristics of
577 palustrine carbonates. (A) The facies of the palustrine carbonates as visible at the scale of the
578 outcrop and on thin section. (B) Reddish Fe-precipitation features along the root-tracks across the
579 laminated sandstones. The black wavy laminae follow Fe-Mn precipitation along sedimentary
580 structures and mark small changes in grain-size. (C) The root tracks on the lower surface marked
581 by stronger cementation. Locally, small voids are still present (indicated by arrows). (D) Remnants
582 of vertical root traces (white arrows) and gastropods (black arrows). (E) Plain view of the upper
583 surface characterized by Fe precipitation and coating around the root-traces.

584 Figure 4. Micromorphological features of palustrine carbonates. (A) Microfacies A-1: packed sands
585 with micritic groundmass made of predominant subangular to angular chert and quartz grains with
586 subordinated rounded to subrounded limestone grains (XPL). (B) Microfacies A-2: groundmass
587 made of dark cyanobacterial clotted micrite (PPL). (C) Microfacies A-2: abundant Fe/Mn
588 precipitation features also with dendrolitic forms (PPL). (D) Microfacies B: sub-horizontal planar
589 voids filled with Fe/Mn features, more frequent toward the top, and impregnation features and
590 masses in the dark cyanobacterial clotted micrite of the groundmass (PPL). (E) Microfacies B:
591 characters of the patches of lighter micritic groundmass with rare microsparitic and sparitic
592 features mainly related to dissolution of shells or as coatings on biological voids (PPL). (F)
593 Microfacies C-1: overall aspect with dark cyanobacterial clotted micrite and very abundant shells,
594 shell fragments and foraminifera (PPL). (G) Microfacies C-2: the alternating patches of dark
595 cyanobacterial clotted micrite and light micrite in the groundmass (PPL). (H) The darker patches

596 of cyanobacterial micrite with abundant microsparitic and sparitic fillings of cracks and biological
597 voids (XPL)

598 Figure 5. Mollusks from the Middle Pleistocene lagoonal mudstones exposed at Torre Mucchia.

599 Vertical bars 2 mm; Horizontal bar 4 mm. (A-B) *Planorbarius corneus*. (C-D) *Theodoxus*
600 *fluviatilis*. (E-F) *Valvata piscinalis*. (G) *Rissoa membranacea*. (H) *Ecrobia ventrosa*. (I)
601 *Cerastoderma glaucum*. (J) *Abra segmentum*.

602 Figure 6. Fish otoliths from the Middle Pleistocene lagoonal mudstones exposed at Torre Mucchia. (A)

603 *Atherina hepsetus* Linnaeus, 1758. (B) *Pomatoschistus* aff. *minutus* (Pallas, 1770).

604 Figure 7. Microphotographs of the main biogenic content of lagoon sandy mudstones. Scale bars 1 mm.

605 (A) Sample 3ORT, vegetal remains, *Ammonia beccarii* (a), limonite nodules (b) and framboidal
606 pyrite (c). (B) Sample 3ORT, *Cribronion granosum* (a), *Valvulineria perlucida* (b) and
607 framboidal pyrite (c). (C) Sample 4ORT, biogenic detritus with *Haynesina paucilocula* (a);
608 *Ammonia beccarii* (b). (D) Sample 4ORT, biogenic detritus with *Ammonia beccarii* (a),
609 *Valvulineria perlucida* (b); *Haynesina paucilocula* (c); *Cyprideis torosa* (d).

610 Figure 8. Pollen percentage diagram of the Torre Mucchia section.

611 Figure 9. Stratigraphic setting of selected Early and Middle Pleistocene pollen records from the Italian
612 Peninsula (42°-39°latitude N), and schematic pollen biostratigraphy of selected pollen taxa
613 (modified from Magri and Palombo, 2013).

614

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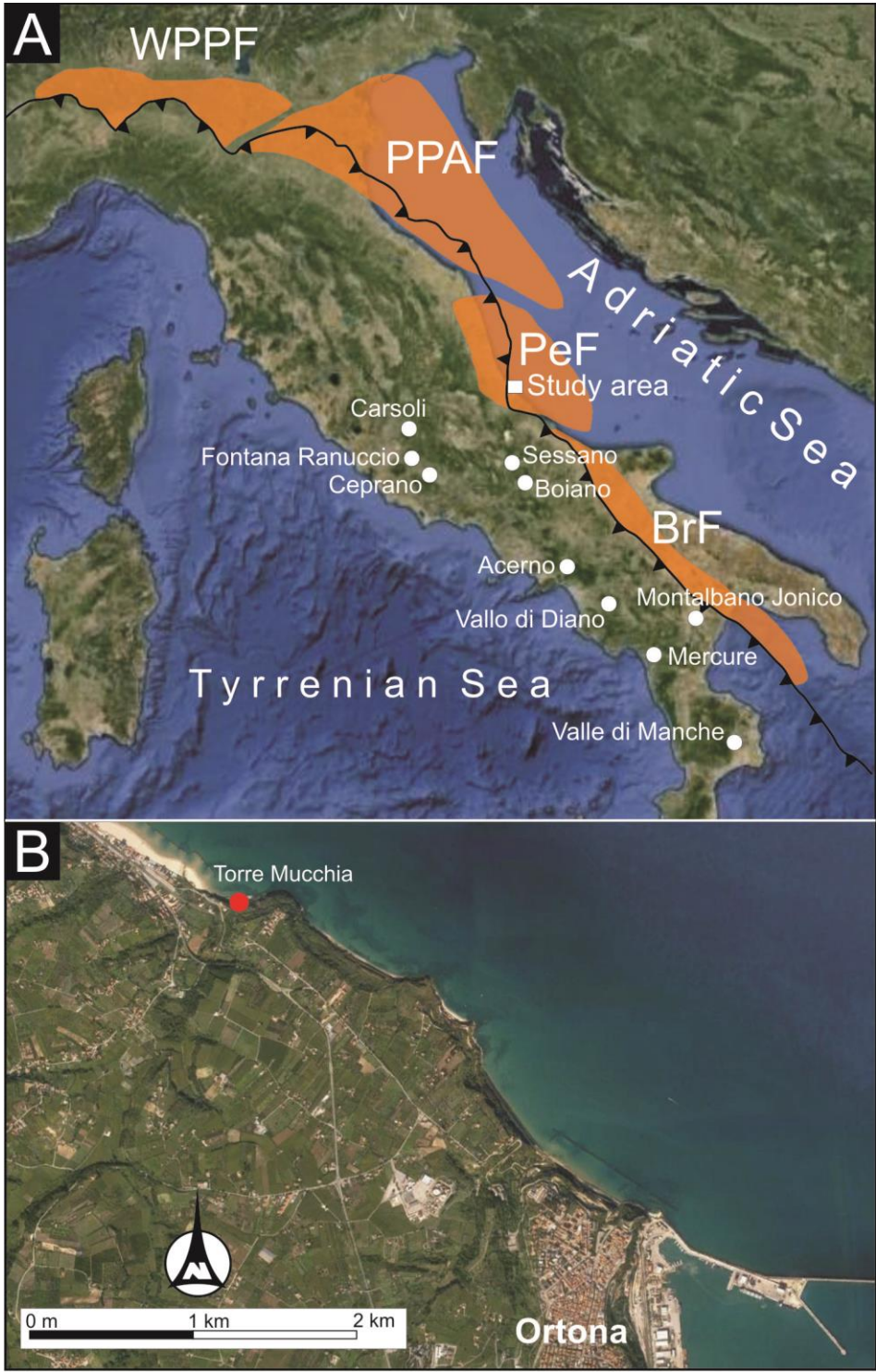


Figure 1

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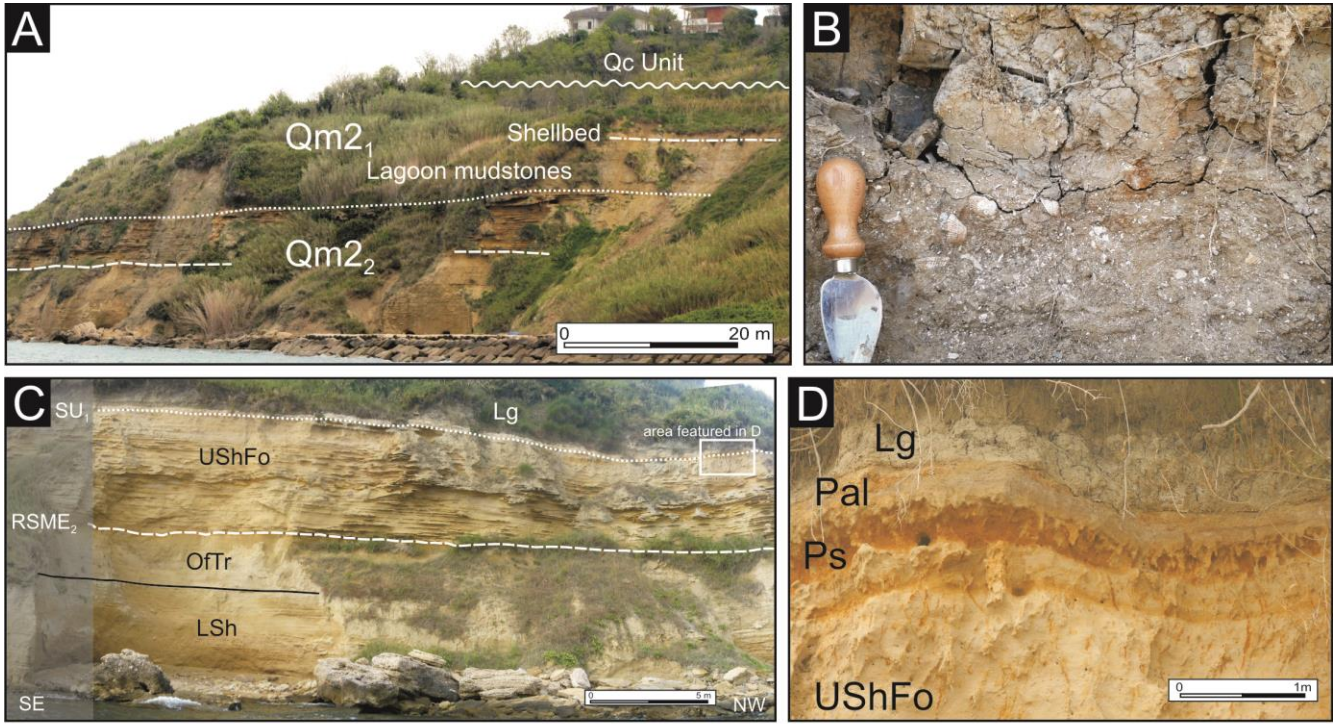


Figure 2

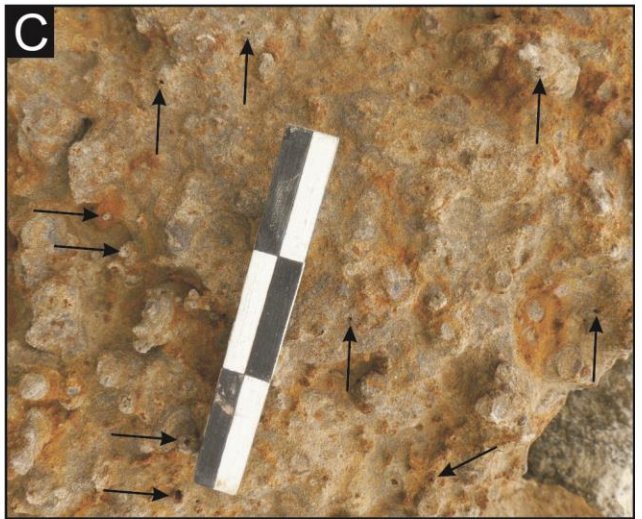
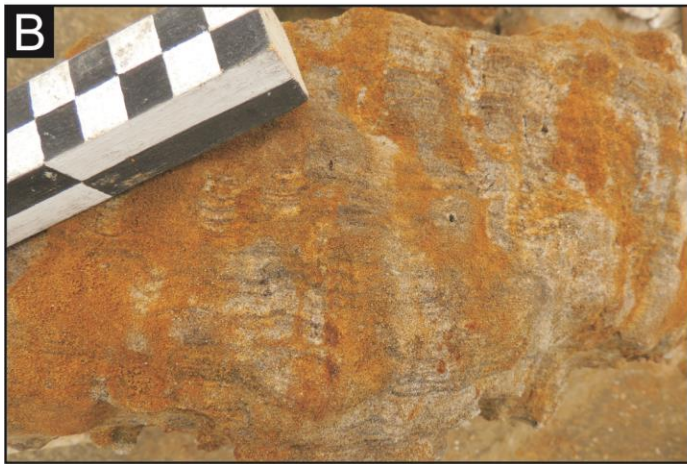
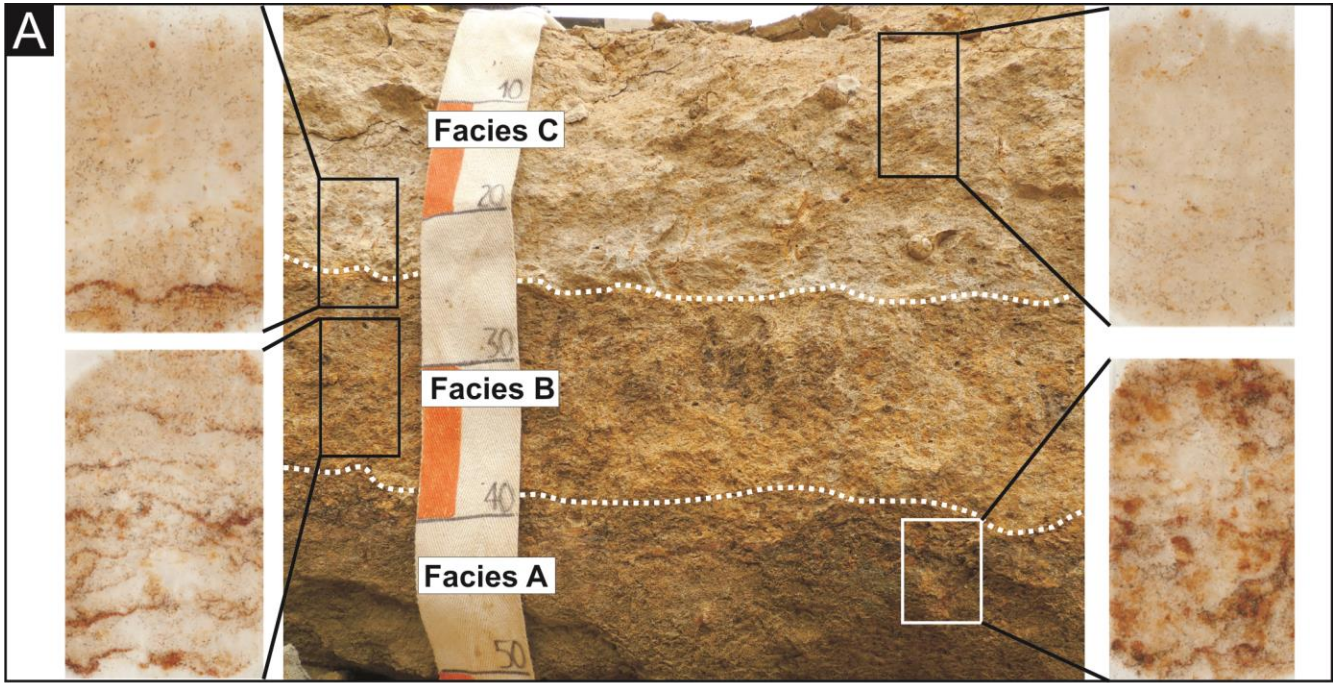
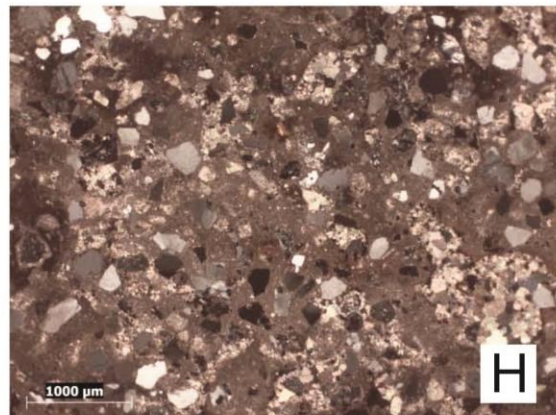
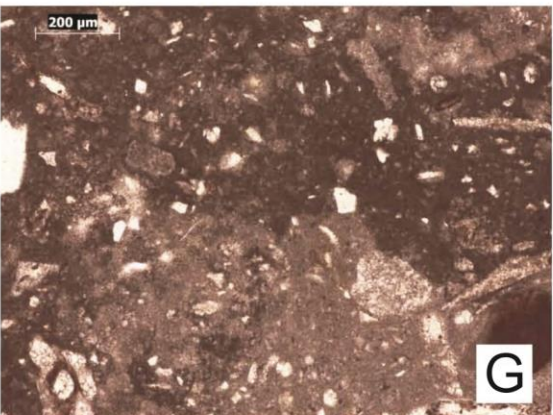
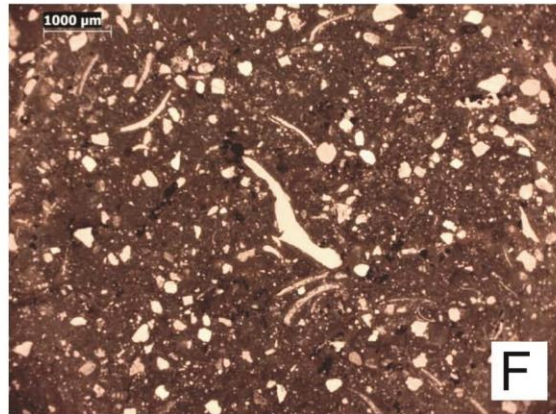
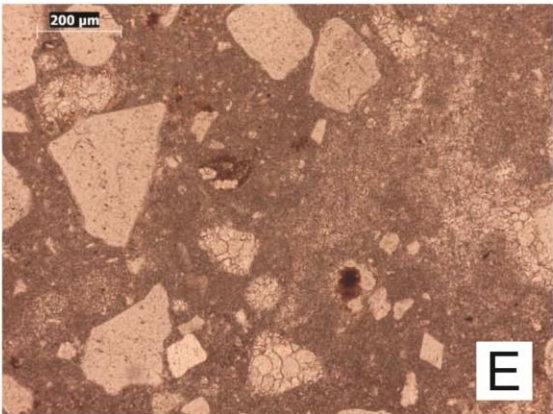
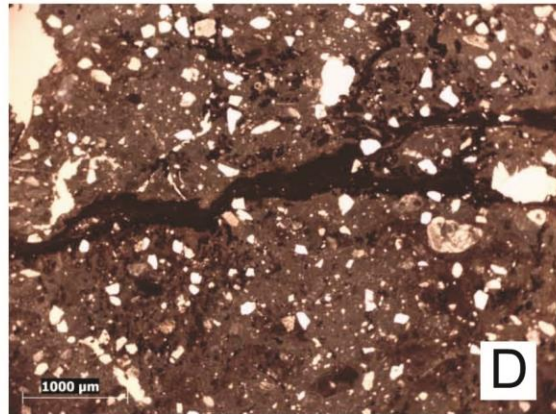
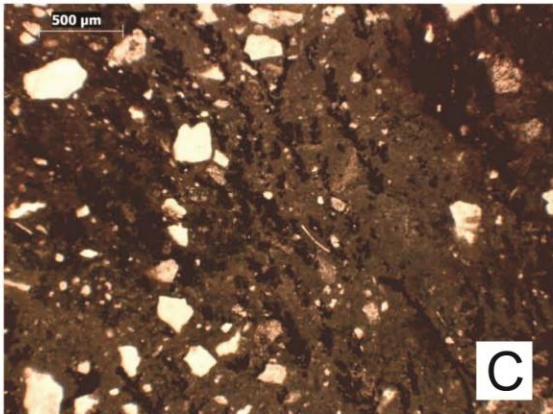
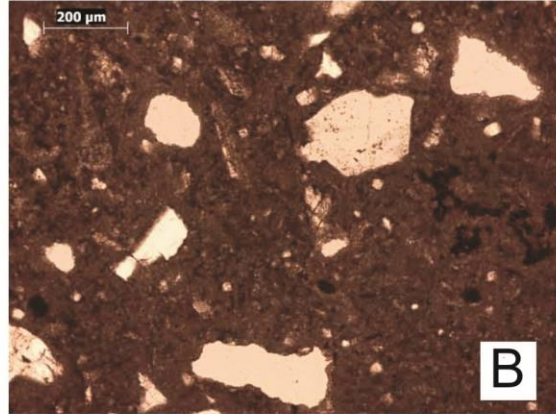
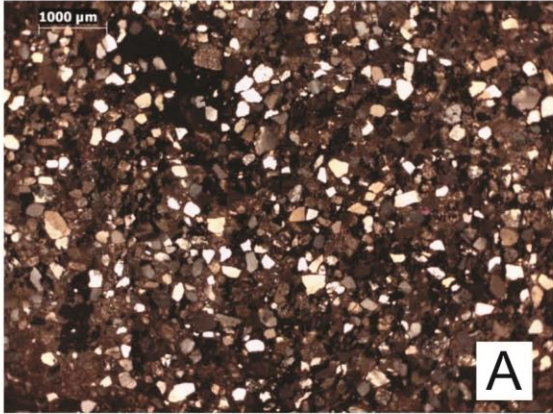
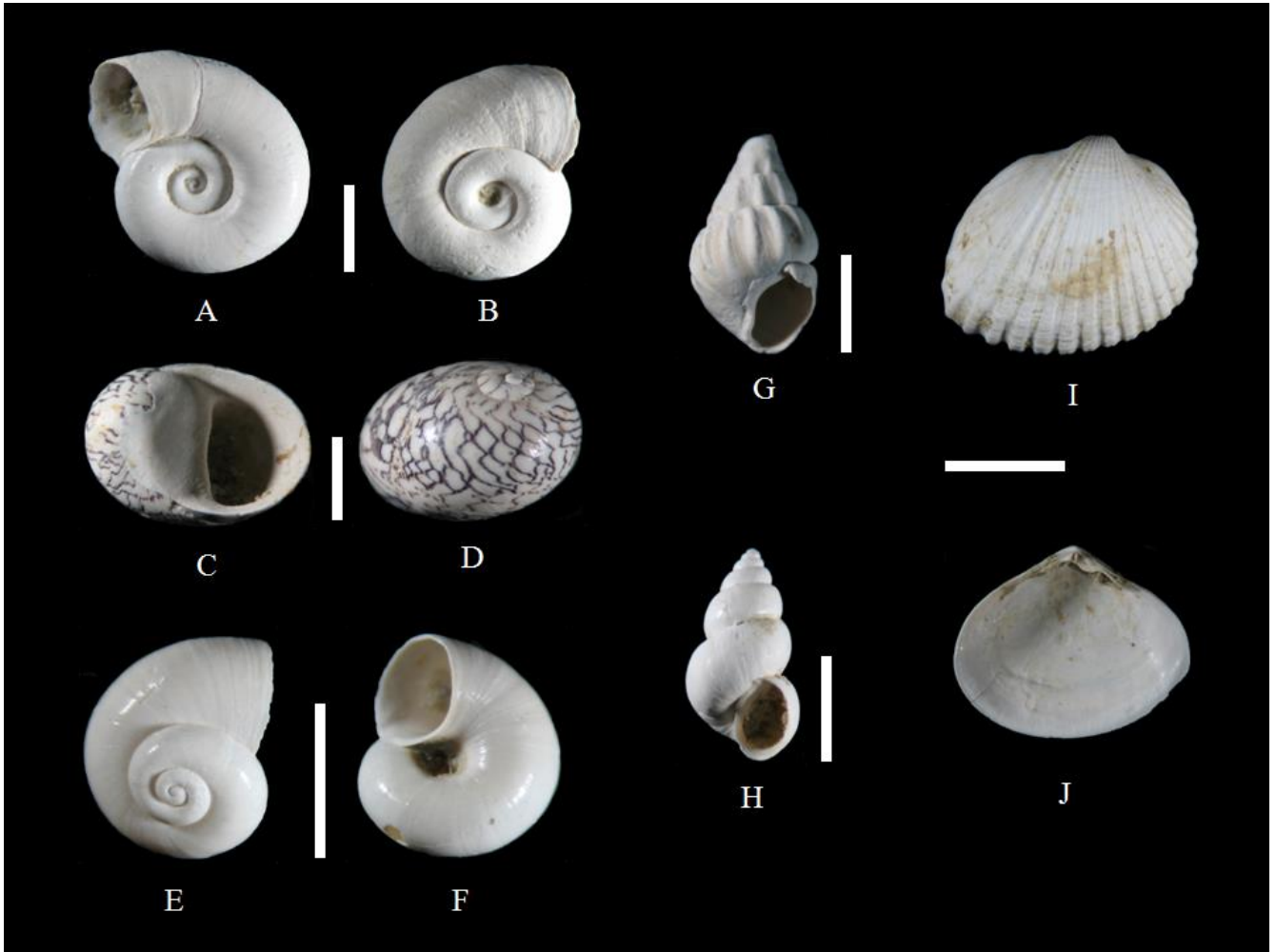


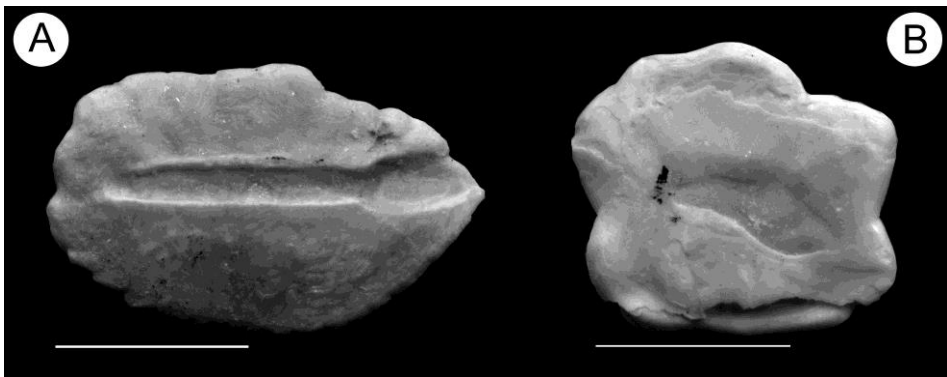
Figure 3





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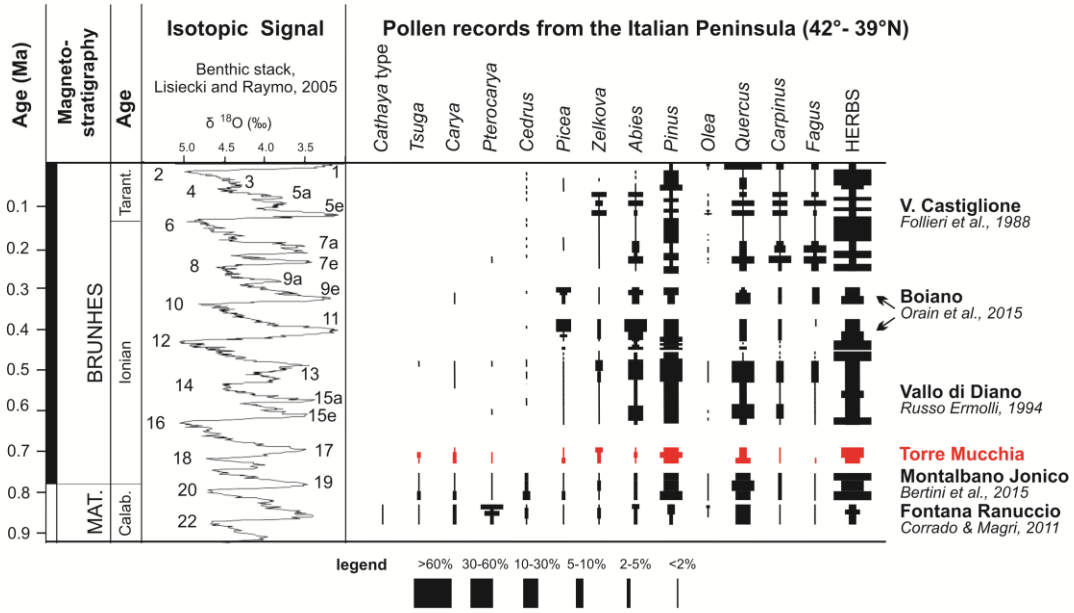
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Supplementary Material

Sedimentology, faunal content and pollen record of Middle Pleistocene palustrine and lagoonal sediments from the Peri-Adriatic basin (Abruzzi, eastern central Italy)

Pierluigi Pieruccini, Claudio Di Celma, Federico Di Rita, Donatella Magri,
Giorgio Carnevale, Piero Farabollini, Luca Ragaini, Mauro Caffau

Torre Muchia pollen counts – Arboreal Pollen

Sample depth (m)	0.6	1	1.2	1.4	3.4	3.6	4	5.2	6	7.6	7.8	8
<i>Abies</i>	2	1	2	0	8	2	1	1		1	1	
<i>Ephedra distachya</i> t.				1					1			
<i>Ephedra fragilis</i> t.	3	6	1	1			3	2	3		1	
<i>Juniperus</i>	3	3	7	1	2	3	4	2	1	1	9	2
<i>Picea</i>	1	6	8	1	4	1	1		5		1	3
<i>Pinus</i>	40	43	31	31	63	21	45	38	44	25	17	20
<i>Acer</i>										1		
<i>Alnus</i>	8	5	9	8	2	1	4	9	5		1	7
<i>Betula</i>	5	5	1		1	3	1		1		8	1
<i>Carpinus betulus</i>	1		1	1		1					5	1
<i>Carya</i>	7	8	6	1		6	2			1	2	1
<i>Corylus</i>	1	1	1	4			4		1		2	2
<i>Ericaceae</i>	1			1		1		2				1
<i>Fagus</i>		1	1									
<i>Fraxinus</i>						1						
<i>Fraxinus ornus</i> t.							1					
<i>Pterocarya</i>				1		1						
<i>Myrtaceae</i>							2					
<i>Carpinus orientalis</i> t.	7	2	1			2	1			4	14	9
<i>Tsuga</i>	1	1			1		4		9			
<i>Pistacia</i>				1								
<i>Quercus deciduous</i>	20	11	5	3	5	5	4	7	1	2	13	5
<i>Quercus cerris/suber</i>											4	
<i>Quercus evergreen</i>	14	10	2	6	4	3	2	5		1	9	7
<i>Ulmus</i>		2		1							1	2
<i>Zelkova</i>	6	7	2	7	3	2	2	11	3	3	4	16